SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANTS: Hiatt, Andrew C.
 Hein, Mich B.
 Fitchen, John H.
- (ii) TITLE OF INVENTION: NOVEL EPITHELIAL TISSUE IMAGING AGENT
- (iii) NUMBER OF SEQUENCES: 93
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: SEED and BERRY LLP
 - (B) STREET: 6300 Columbia Center, 701 Fifth Avenue
 - (C) CITY: Seattle
 - (D) STATE: Washington
 - (E) COUNTRY: USA
 - (F) ZIP: 98104
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE: 9-JAN-1998
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Maki, David J.
 - (B) REGISTRATION NUMBER: 31,392
 - (C) REFERENCE/DOCKET NUMBER: 310098.402
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (206) 622-4900
 - (B) TELEFAX: (206) 682-6031
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 137 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Gln Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys Cys Lys Cys Ala 1 5 10 15

Arg Ile Thr Ser Arg Ile Ile Arg Ser Ser Glu Asp Pro Asn Glu Asp 20 25 30

Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg Glu 35 40 45

Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Arg Pro Val Tyr His 50 55 60

Leu Ser Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu Asp

70

75

80

Asn Gln Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp Ser 85 90 95

Ala Thr Glu Thr Cys Tyr Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Ala 100 105 110

Val Val Pro Leu Val Tyr Gly Gly Glu Thr Lys Met Val Glu Thr Ala 115 120 125

Leu Thr Pro Asp Ala Cys Tyr Pro Asp 130

- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 135 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Gln Asp Glu Asn Glu Arg Ile Val Val Asp Asn Lys Cys Lys Cys Ala 1 5 10 15

Arg Ile Thr Ser Arg Ile Ile Pro Ser Ala Glu Asp Pro Ser Gln Asp 20 25 30

Ile Val Glu Arg Asn Val Arg Ile Ile Val Pro Leu Asn Ser Arg Glu 35 40 45

Asn Ile Ser Asp Pro Thr Ser Pro Met Arg Thr Lys Pro Val Tyr His 50 55 60

Leu Ser Asp Leu Cys Lys Lys Cys Asp Thr Thr Glu Val Glu Leu Glu 65 70 75 80

Asp Gln Val Val Thr Ala Ser Gln Ser Asn Ile Cys Asp Ser Asp Ala 85 90 95

Glu Thr Cys Tyr Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Asn Arg Val 100 105 110

Lys Leu Ser Tyr Arg Gly Gln Thr Lys Met Val Glu Thr Ala Leu Thr
115 120 125

Pro Asp Ser Cys Tyr Pro Asp 130 135

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 137 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Asp Asp Glu Ala Thr Ile Leu Ala Asp Asn Lys Cys Met Cys Thr Arg 1 5 10 15

Val Thr Ser Arg Ile Ile Pro Ser Thr Glu Asp Pro Asn Glu Asp Ile 20 25 30

Val Glu Arg Asn Ile Arg Ile Val Val Pro Leu Asn Asn Arg Glu Asn 35 40 45

Ile Ser Asp Pro Thr Ser Pro Leu Arg Arg Asn Pro Val Tyr His Leu 50 55 60

Ser Asp Val Cys Lys Lys Cys Asp Pro Val Glu Val Glu Leu Glu Asp 65 70 75 80

Gln Val Val Thr Ala Thr Gln Ser Asn Ile Cys Asn Glu Asp Asp Gly 85 90 95

Val Pro Glu Thr Cys Tyr Met Tyr Asp Arg Asn Lys Cys Tyr Thr Thr 100 105 110

Met Val Pro Leu Arg Tyr His Gly Glu Thr Lys Met Val Gln Ala Ala 115 120 125 Leu Thr Pro Asp Ser Cys Tyr Pro Asp 130 135

- (2) INFORMATION FOR SEQ ID NO:4:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 136 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Glu Asp Glu Ser Thr Val Leu Val Asp Asn Lys Cys Gln Cys Val Arg 1 5 10 15

Ile Thr Ser Arg Ile Ile Arg Asp Pro Asp Asn Pro Ser Glu Asp Ile
20 25 30

Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Thr Arg Glu Asn 35 40 45

Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Glu Pro Lys Tyr Asn Leu 50 55 60

Ala Asn Leu Cys Lys Lys Cys Asp Pro Thr Glu Ile Glu Leu Asp Asn 65 70 75 80

Gln Val Phe Thr Ala Ser Gln Ser Asn Ile Cys Pro Asp Asp Asp Tyr 85 90 95

Ser Glu Thr Cys Tyr Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Thr Leu 100 105 110

Val Pro Ile Thr His Arg Gly Val Thr Arg Met Val Lys Ala Thr Leu 115 120 125

Thr Pro Asp Ser Cys Tyr Pro Asp 130 135

- (2) INFORMATION FOR SEQ ID NO:5:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 119 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Glu Gln Glu Tyr Ile Leu Ala Asn Asn Lys Cys Lys Cys Val Lys Ile 1 5 10 15

Ser Ser Arg Phe Val Pro Ser Thr Glu Arg Pro Gly Glu Glu Ile Leu 20 25 30

Glu Arg Asn Ile Gln Ile Thr Ile Pro Thr Ser Ser Arg Met Xaa Ile 35 40 45

Ser Asp Pro Tyr Ser Pro Leu Arg Thr Gln Pro Val Tyr Asn Leu Trp 50 55 60

Asp Ile Cys Gln Lys Cys Asp Pro Val Gln Leu Glu Ile Gly Gly Ile 65 70 75 80

Pro Val Leu Ala Ser Gln Pro Xaa Xaa Ser Xaa Pro Asp Asp Glu Cys 85 90 95

Tyr Thr Thr Glu Val Asn Phe Lys Lys Lys Val Pro Leu Thr Pro Asp 100 105 110

Ser Cys Tyr Glu Tyr Ser Glu 115

(2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 128 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Asn Lys Cys Met Cys Thr Arg Val Thr Ala Arg Ile Arg Gly Thr Arg 1 5 10 15

Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Tyr Ile Arg Ile Asn Val 20 25 30

Pro Leu Lys Asn Arg Gly Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg 35 40 45

Asn Gln Pro Val Tyr His Leu Ser Pro Ser Cys Lys Lys Cys Asp Pro 50 55 60

Tyr Glu Asp Gly Val Val Thr Ala Thr Glu Thr Asn Ile Cys Tyr Pro
65 70 75 80

Asp Gln Gly Val Pro Gln Ser Cys Arg Asp Tyr Cys Pro Glu Leu Asp 85 90 95

Arg Asn Lys Cys Tyr Thr Val Leu Val Pro Pro Gly Tyr Thr Gly Glu
100 105 110

Thr Lys Met Val Gln Asn Ala Leu Thr Pro Asp Ala Cys Tyr Pro Asp 115 120 125

(2) INFORMATION FOR SEQ ID NO:7:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 421 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..414

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

											AAC Asn						48
											GAG Glu						96
											CCA Pro					Ξ	144
											ACA Thr 60					-	192
											ACA Thr					2	240
											ATT Ile					2	288
AGC	GCT	ACA	GAA	ACC	TGC	AGC	ACC	TAC	GAT	AGG	AAC	AAA	TGC	TAC	ACG	3	336

Ser	Ala	Thr	Glu 100	Thr	Cys	Ser	Thr	Tyr 105	Asp	Arg	Asn	Lys	Cys 110	Tyr	Thr	
GCC Ala	GTG Val	GTT Val 115	CCG Pro	CTC Leu	GTG Val	TAT Tyr	GGT Gly 120	GGA Gly	GAG Glu	ACA Thr	AAA Lys	ATG Met 125	GTG Val	GAA Glu	ACT Thr	384
GCC Ala	CTT Leu 130	ACG Thr	CCC Pro	GAT Asp	GCA Ala	TGC Cys 135	TAT Tyr	CCG Pro	GAC Asp	TGA	ATTC					421
(2)) SE(QUENCA) LI B) T'	FOR CE CI ENGTI YPE: TRANI	HARAG H: 2: nucl	CTER 15 ba leic ESS:	ISTIC ase p acic sing	CS: pair:	5							
		(I	A) N2	E: AME/I DCATI	ON:	12		SEQ I	ID NO	D:8:						
GAT Asp 1	CAG Gln	AAG Lys	TGC Cys	AAG Lys 5	TGT Cys	GCT Ala	CGT Arg	ATT Ile	ACT Thr 10	TCT Ser	AGA Arg	ATC Ile	ATC Ile	CGT Arg 15	AGC Ser	48
TCA Ser	GAG Glu	GAC Asp	CCA Pro 20	AAT Asn	GAA Glu	GAT Asp	ATA Ile	GTC Val 25	GAA Glu	CGT Arg	AAC Asn	ATC Ile	CGT Arg 30	ATC Ile	ATC Ile	96
GTC Val	CCA Pro	CTG Leu 35	AAT Asn	AAC Asn	CGG Arg	GAG Glu	AAT Asn 40	ATC Ile	TCA Ser	GAT Asp	CCT Pro	ACA Thr 45	AGT Ser	CCG Pro	TTG Leu	144
CGC Arg	ACA Thr 50	CGC Arg	TTC Phe	GTA Val	TAC Tyr	CAC His 55	CTG Leu	TCA Ser	GAT Asp	CTG Leu	TGT Cys 60	AAG Lys	AAG Lys	GAT Asp	GAG Glu	192
				GAA Glu			TG									215
(2)	INFO	RMAT	'ION	FOR	SEQ	ID N	0:9:									

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 140 base pairs(B) TYPE: nucleic acid

(C) STRANDEDNESS: single(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
CTAGAATCAT CCGTAGCTCA GAGGACCCAA ATGAAGATAT AGTCGAACGT AACATCCGTA	60
TCATCGTCCC ACTGAATAAC CGGGAGAATA TCTCAGATCC TACAAGTCCG TTGCGCACAC	120
GCTTCGTATA CCACCTGTCA	140
(2) INFORMATION FOR SEQ ID NO:10:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: GATCAGAAGT GCAAGTGTGC TCGTATTACT T (2) INFORMATION FOR SEQ ID NO:11:	31
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 44 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 142	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
GAT CTG TGT AAG AAG GAT GAA GAT TCC GCT ACA GAA ACC TGC Asp Leu Cys Lys Lys Asp Glu Asp Ser Ala Thr Glu Thr Cys 75 80 85	42
rg	11

(2)) IN	FORM	ATIC	N FO	R SE	Q ID	NO:	12:								
	(i	((A) I B) T C) S	CE C ENGT YPE: TRAN	H: 1 nuc DEDN	09 b leic ESS:	ase aci sin	pair d	S							
	, ,	٠														
~~-				CE D												
														GGTG	GAGAGA	k 60
				ACTG					GCAT	GCTA	CCC	TGAC'	TG			109
(2)	INF	ORMA'	TION	FOR	SEQ	ID :	NO:1	3:								
	(i)	() ()	A) L B) T C) S	CE C ENGT YPE: TRAN	H: 2 nuc DEDN	86 b leic ESS:	ase aci sin	pair d	S							
		() (1	B) L	AME/I	ION:	1										
~~~				CE DI												
GAC Asp 15	AAC Asn	AAG Lys	TGC Cys	AAG Lys	TGT Cys 20	GCT Ala	CGT Arg	ATT Ile	ACT Thr	TCT Ser 25	AGA Arg	ATC Ile	ATC Ile	CGT Arg	AGC Ser 30	48
TCA	GAG Glu	GAC Asp	CCA	AAT	GAA	GAT	ATA	GTC	GAA	CGT	AAC	ATC	CGT	ATC	ATC	96
Ser		1	FIO	Asn 35	GIU	Asp	TTE	Val	Glu 40	Arg	Asn	Ile	Arg	Ile 45	Ile	
Ser GTC Val	CCA	CTG	AAT	35 AAC	CGG	GAG	AAT	ATC	40 TCA	GAT	CCT	ACA	AGT	45 CCG	TTG	144
GTC	CCA Pro ACA	CTG Leu CGC	AAT Asn 50 TTC	35 AAC Asn GTA	CGG Arg TAC	GAG Glu CAC	AAT Asn CTG	ATC Ile 55 TCA	40 TCA Ser	GAT Asp CTG	CCT Pro	ACA Thr	AGT Ser 60	45 CCG Pro	TTG Leu GAT	144 192

AAC ATT TGC GAT GAG GAC AGC GCT ACA GAA ACC TGC TAC TGA Asn Ile Cys Asp Glu Asp Ser Ala Thr Glu Thr Cys Tyr * 95 100 105	282
ATTC	286
(2) INFORMATION FOR SEQ ID NO:14:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 105 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
<pre>(ix) FEATURE:     (A) NAME/KEY: CDS     (B) LOCATION: 1105  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:</pre>	
GAT CTG TGT AAG AAG TGT GAT CCA ACA GAG GTA GAG CTG GAC AAT CAG	48
Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu Asp Asn Gln 95 100 105 110	
ATA GTC ACT GCG ACT CAA AGC AAC ATT TGC GAT GAG GAC AGC GCT ACA Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp Ser Ala Thr 115 120 125	96
CTT TGG ACG Leu Trp Thr	105
(2) INFORMATION FOR SEQ ID NO:15:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 61 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
GATCAGGAAG ATGAACGTAT TGTTCTGGTT GACAACAAGT GCAAGTGTGC TCGTATTACT	60
т	61

(2)	INF	'ORMA	TION	f FOR	SEQ	ID	NO:1	.6:							
	(i	( (	A) L B) T C) S	CE C ENGT YPE: TRAN	H: 1 nuc DEDN	98 b leic ESS:	ase aci sin	pair d	S						
	(xi	) SE	QUEN	CE D	ESCR	IPTI	ON:	SEQ	ID N	0:16	:				
GCG.	ATGA	CGA	CGAT.	AAGG	CC C.	AAAC	GGAG	A CC	TGTA	CTGT	TGC	GCCT	CGT	GAAC	GGCAAA
ACT	GCGG.	ATT	CCCG	GAAG'	TA A	CACC	CTCT	C AG	TGCG	CTAA	TAA	AGGC	TGC	TGTT	TTGATG
ACA	CGGT.	ACG	GGGC	GTTC	CG T	GGTG	CTTC	T AC	CCCA	ATAC	AAT	TGAC	GTT	CCGC	CTGAAG
AAG	AGTG	CGA	GCCG'	TAAG											
(2)	INF	ORMA	TION	FOR	SEQ	ID 1	NO:1	7:							
		(i)	(A (B		NGTH PE: a	: 13: amin	8 am			s					
	(:	ii) 1	MOLE	CULE	TYPI	E: p	rote	in							
	(:	xi) :	SEQU!	ENCE	DES	CRIP:	TION	: SE	Q ID	NO:	17:				
Asp 1	Gln	Glu	Asp	Glu 5	Arg	Ile	Val	Leu	Val 10	Asp	Asn	Lys	Cys	Lys 15	Cys
Ala	Arg	Ile	Thr 20	Ser	Arg	Ile	Ile	Arg 25	Ser	Ser	Glu	Asp	Pro 30	Asn	Glu
Asp	Ile	Val 35	Glu	Arg	Asn	Ile	Arg 40	Ile	Ile	Val	Pro	Leu 45	Asn	Asn	Arg
Glu	Asn 50	Ile	Ser	Asp	Pro	Thr 55	Ser	Pro	Leu	Arg	Thr 60	Arg	Phe	Val	Tyr
His 65	Leu	Ser	Asp	Leu	Cys 70	Lys	Lys	Cys	Asp	Pro 75	Thr	Glu	Val	Glu	Leu 80
Asp	Asn	Gln	Ile	Val 85	Thr	Ala	Thr	Gln	Ser 90	Asn	Ile	Cys	Asp	Glu 95	Asp

Ser Ala Thr Glu Thr Cys Ser Thr Tyr Asp Arg Asn Lys Cys Tyr Thr 100 105 110

Ala Val Val Pro Leu Val Tyr Gly Gly Glu Thr Lys Met Val Glu Thr 115 120 125

Ala Leu Thr Pro Asp Ala Cys Tyr Pro Asp 130 135

- (2) INFORMATION FOR SEQ ID NO:18:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 71 amino acids
    - (B) TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: protein
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Asp Gln Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser 1 5 10 15

Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile 20 25 30

Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu
35 40 45

Arg Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Asp Glu 50 55 60

Asp Ser Ala Thr Glu Thr Cys 65 70

- (2) INFORMATION FOR SEQ ID NO:19:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 49 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Ser Arg Ile Ile Arg Ser Ser Glu Asp Pro Asn Glu Asp Ile Val Glu 1 5 10 15

Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg Glu Asn Ile Ser 20 25 30

Asp Pro Thr Ser Pro Leu Arg Thr Arg Phe Val Tyr His Leu Ser Asp 35 40 45

Leu

- (2) INFORMATION FOR SEQ ID NO:20:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 12 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

Asp Gln Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg 1 5 10

- (2) INFORMATION FOR SEQ ID NO:21:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 14 amino acids
    - (B) TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: protein
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

Asp Leu Cys Lys Lys Asp Glu Asp Ser Ala Thr Glu Thr Cys

1 5 10

- (2) INFORMATION FOR SEQ ID NO:22:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 36 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

Ser Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Ala Val Val Pro Leu Val 1 5 10 15

Tyr Gly Glu Thr Lys Met Val Glu Thr Ala Leu Thr Pro Asp Ala 20 25 30

Cys Tyr Pro Asp 35

- (2) INFORMATION FOR SEQ ID NO:23:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 93 amino acids
    - (B) TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: protein
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

Asp Asn Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser

1 5 10 15

Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile 20 25 30

Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu 35 40 45

Arg Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Cys Asp 50 55 60

Pro Thr Glu Val Glu Leu Asp Asn Gln Ile Val Thr Ala Thr Gln Ser 65 70 75 80

Asn Ile Cys Asp Glu Asp Ser Ala Thr Glu Thr Cys Tyr 85 90

- (2) INFORMATION FOR SEQ ID NO:24:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 35 amino acids
    - (B) TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: protein
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu Asp Asn Gln
1 5 10 15

Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp Ser Ala Thr
20 25 30

Leu Trp Thr

35

- (2) INFORMATION FOR SEQ ID NO:25:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 22 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

Asp Gln Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys Cys Lys Cys 1 5 10 15

Ala Arg Ile Thr Ser Arg
20

- (2) INFORMATION FOR SEQ ID NO:26:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 66 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

Cys Ser Asp Asp Asp Asp Lys Ala Gln Thr Glu Thr Cys Thr Val Ala 1 5 10 15

Pro Arg Glu Arg Gln Asn Cys Gly Phe Pro Gly Val Thr Pro Ser Gln 20 25 30

Cys Ala Asn Lys Gly Cys Cys Phe Asp Asp Thr Val Arg Gly Val Pro
35 40 45

Trp Cys Phe Tyr Pro Asn Thr Ile Asp Val Pro Pro Glu Glu Glu Cys 50 55 60

Glu Phe

- (2) INFORMATION FOR SEQ ID NO:27:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 421 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single

### (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2	7 :
----------------------------------------	-----

CTAGTCCTTC	TACTTGCATA	ACAAGACCAA	CTGTTGTTCA	CGTTCACACG	AGCATAATGA	60
AGATCTTAGT	AGGCATCGAG	TCTCCTGGGT	TTACTTCTAT	ATCAGCTTGC	ATTGTAGGCA	120
TAGTAGCAGG	GTGACTTATT	GGCCCTCTTA	TAGAGTCTAG	GATGTTCAGG	CAACGCGTGT	180
GCGAAGCATA	TGGTGGACAG	TCTAGACACA	TTCTTCACAC	TAGGTTGTCT	CCATCTCGAC	240
CTGTTAGTCT	ATCAGTGACG	CTGAGTTTCG	TTGTAAACGC	TACTCCTGTC	GCGATGTCTT	300
TGGACGTCGT	GGATGCTATC	CTTGTTTACG	ATGTGCCGGC	ACCAAGGCGA	GCACATACCA	360
CCTCTCTGTT	TTTACCACCT	TTGACGGGAA	TGCGGGCTAC	GTACGATAGG	CCTGACTTAA	420
G						421

# (2) INFORMATION FOR SEQ ID NO:28:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 219 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

# (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

CTAGTCTTCA CGTTCACACG AGCATAATGA AGATCTTAGT AGGCATCGAG TCTCCTGGGT 60

TTACTTCTAT ATCAGCTTGC ATTGTAGGCA TAGTAGCAGG GTGACTTATT GGCCCTCTTA 120

TAGAGTCTAG GATGTTCAGG CAACGCGTGT GCGAAGCATA TGGTGGACAG TCTAGACACA 180

TTCTTCCTAC TCCTGTCGCG ATGTCTTGG ACGACTTAA 219

# (2) INFORMATION FOR SEQ ID NO:29:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 140 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:	
TTAGTAGGCA TCGAGTCTCC TGGGTTTACT TCTATATCAG CTTGCATTGT AGGCATAGTA	60
GCAGGGTGAC TTATTGGCCC TCTTATAGAG TCTAGGATGT TCAGGCAACG CGTGTGCGAA	120
GCATATGGTG GACAGTCTAG	140
(2) INFORMATION FOR SEQ ID NO:30:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 31 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:	
TCTTCACGTT CACACGAGCA TAATGAAGAT C	31
(2) INFORMATION FOR SEQ ID NO:31:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 44 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31: ACACATTCTT CCTACTTCTC AGGCGATGTC TTTGGACGAC TTAA	
(2) INFORMATION FOR SEQ ID NO:32:	44
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 117 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	

(X1) SEQUENCE DESCRIPTION: SEQ ID NO:32:	
ACGTCGTGGA TGCTATCCTT GTTTACGATG TGCCGGCACC AAGGCGAGCA CATACCACCT	60
CTCTGTTTTT ACCACCTTTG ACGGGAATGC GGGCTACGTA CGATGGGACT GACTTAA	117
(2) INFORMATION FOR SEQ ID NO:33:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 282 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:	
CTGTTGTTCA CGTTCACACG AGCATAATGA AGATCTTAGT AGGCATCGAG TCTCCTGGGT	60
TTACTTCTAT ATCAGCTTGC ATTGTAGGCA TAGTAGCAGG GTGACTTATT GGCCCTCTTA	120
TAGAGTCTAG GATGTTCAGG CAACGCGTGT GCGAAGCATA TGGTGGACAG TCTAGACACA	180
TTCTTCACAC TAGGTTGTCT CCATCTCGAC CTGTTAGTCT ATCAGTGACG CTGAGTTTCG	240
TTGTAAACGC TACTCCTGTC GCGATGTCTT TGGACGATGA CT	282
(2) INFORMATION FOR SEQ ID NO:34:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 105 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:	
GATCTGTGTA AGAAGTGTGA TCCAACAGAG GTAGAGCTGG ACAATCAGAT AGTCACTGCG	60
ACTCAAAGCA ACATTTGCGA TGAGGACAGC GCTACACTTT GGACG	105
(2) INFORMATION FOR SEQ ID NO:35:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 65 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:
CTAGTCCTTC TACTTGCATA ACAAGACCAA CTGTTGTTCA CGTTCACACG AGCATAATGA 60
AGATC 69
(2) INFORMATION FOR SEQ ID NO:36:
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 206 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:
ACTTCGCTAC TGCTGCTATT CCGGGTTTGC CTCTGGACAT GACAACGCGG AGCACTTGCC 60
GTTTTGACGC CTAAGGGCCT TCATTGTGGG AGAGTCACGC GATTATTTCC GACGACAAAA 120
CTACTGTGCC ATGCCCCGCA AGGCACCACG AAGATGGGGT TATGTTAACT GCAAGGCGGA 180
CTTCTTCTCA CGCTCGGCAT TCTTAA 206
(2) INFORMATION FOR SEQ ID NO:37:
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 13 amino acids</li><li>(B) TYPE: amino acid</li><li>(C) STRANDEDNESS:</li><li>(D) TOPOLOGY: linear</li></ul>

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:

Asp Gln Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys 1 5 10

- (2) INFORMATION FOR SEQ ID NO:38:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 7 amino acids
    - (B) TYPE: amino acid

- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

Glu Asn Leu Tyr Phe Gln Ser

- (2) INFORMATION FOR SEQ ID NO:39:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 11 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:

Lys Ala His Lys Val Asp Met Val Gln Tyr Thr 1 5 10

- (2) INFORMATION FOR SEQ ID NO:40:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 4 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:

Val Gln Tyr Thr 1

.

- (2) INFORMATION FOR SEQ ID NO:41:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 6 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:

Glu Lys Ala Val Ala Asp 1 5	
(2) INFORMATION FOR SEQ ID NO:42:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 131 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(ix) FEATURE:  (A) NAME/KEY: CDS  (B) LOCATION: 178	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:	
ATG AAA TTC TTA GTC AAC GTT GCC CTT TTT ATG GTC GTA TAC ATT TCT Met Lys Phe Leu Val Asn Val Ala Leu Phe Met Val Val Tyr Ile Ser 40 45 50	48
TAC ATC TAT GCG GAT CCG AGC TCG AGT GCT CTAGATCTGC AGCTGGTACC Tyr Ile Tyr Ala Asp Pro Ser Ser Ala 55 60	98
ATGGAATTCG AAGCTTGGAG TCGACTCTGC TGA	131
(2) INFORMATION FOR SEQ ID NO:43:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 26 amino acids</li><li>(B) TYPE: amino acid</li><li>(D) TOPOLOGY: linear</li></ul>	
(ii) MOLECULE TYPE: protein	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:	
Met Lys Phe Leu Val Asn Val Ala Leu Phe Met Val Val Tyr Ile Ser 1 5 10 15	
Tyr Ile Tyr Ala Asp Pro Ser Ser Ala 20 25	
(2) INFORMATION FOR SEQ ID NO:44:	

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 4 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS:
  - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:

Lys Asp Glu Leu

- (2) INFORMATION FOR SEQ ID NO:45:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 16 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:

Ala Ile Gln Asp Pro Arg Leu Phe Ala Glu Glu Lys Ala Val Ala Asp

- (2) INFORMATION FOR SEQ ID NO:46:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 61 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:

GATCAGGAAG ATGAACGTAT TGTTCTGGTT GACAACAAGT GCAAGTGTGC TCGTATTACT

61

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- (2) INFORMATION FOR SEQ ID NO:47:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 61 base pairs

(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:	
CTAGAAGTAA TACGAGCACA CTTGCACTTG TTGTCAACCA GAACAATACG TTCATCTTCC	60
${f T}$	61
(2) INFORMATION FOR SEQ ID NO:48:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 31 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:48:	
GATCAGAAGT GCAAGTGTGC TCGTATTACT T	31
(2) INFORMATION FOR SEQ ID NO:49:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 31 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:	
CTAGAAGTAA TACGAGCACA CTTGCACTTC T	31
(2) INFORMATION FOR SEQ ID NO:50:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 61 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:	
GATCAGGAAG ATGAACGTAT TGTTCTGGTT GACAACAAGT GCAAGTCCGC TCGTATTACT	50
T	51
(2) INFORMATION FOR SEQ ID NO:51:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 61 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(vi) SEQUENCE DESCRIPTION, SEQ ID NO.51.	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:51:	
CTAGAAGTAA TACGAGCGGA CTTGCACTTG TTGTCAACCA GAACAATACG TTCATCTTCC	50
T	51
(2) INFORMATION FOR SEQ ID NO:52:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 61 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:52:	
GATCAGGAAG ATGAACGTAT TGTTCTGGTT GACAACAAGT GCAAGGTTGC TCGTATTACT	60
Т	61
(2) INFORMATION FOR SEQ ID NO:53:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 61 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	

(xi) SEQUENCE DESC	RIPTION: SEQ ID NO:53:		
CTAGAAGTAA TACGAGCAAC	CTTGCACTTG TTGTCAACCA	GAACAATACG TTCATC	TTCC 60
Т			61
(2) INFORMATION FOR SEC	Q ID NO:54:		
(i) SEQUENCE CHARA  (A) LENGTH: (B) TYPE: nuc  (C) STRANDEDI  (D) TOPOLOGY	17 base pairs cleic acid NESS: single		
(xi) SEQUENCE DESC	RIPTION: SEQ ID NO:54:		
CTAGAATCAT CCGTAGCTCA	GAGGACCCAA ATGAAGATAT	AGTCGAA	47
(2) INFORMATION FOR SE	Q ID NO:55:		
(i) SEQUENCE CHAR.  (A) LENGTH:  (B) TYPE: nu.  (C) STRANDED.  (D) TOPOLOGY	58 base pairs cleic acid NESS: single		
(xi) SEQUENCE DESC	RIPTION: SEQ ID NO:55:		
GATACGGATG TTACGTTCGA	CTATATCTTC ATTTGGGTCC	TCTGAGCTAC GGATGA	ATT 58
(2) INFORMATION FOR SE	Q ID NO:56:		
(i) SEQUENCE CHAR  (A) LENGTH:  (B) TYPE: nu  (C) STRANDED  (D) TOPOLOGY	49 base pairs cleic acid NESS: single		

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:

CGTAACATCC GTATCATCGT CCCACTGAAT AACCGGGAGA ATATCTCAG

(2) INFORMATION FOR SEQ ID NO:57:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 49 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:57:	
CGTAACATCC GTATCATCGT CCCACTGAAT AACCGGGAGC ACATCTCAG	49
(2) INFORMATION FOR SEQ ID NO:58:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 49 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:58:  ACGGACTTGT AGGATCTGAG ATATTCTCCC GGTTATTCAG TGGGACGAT	49
(2) INFORMATION FOR SEQ ID NO:59:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 49 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:59:	
ACGGACTTGT AGGATCTGAG ATGTGCTCCC GGTTATTCAG TGGGACGAT	4.9
(2) INFORMATION FOR SEQ ID NO:60:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 44 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single	

(D) TOPOLOGY: linear

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 44 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:60:	
ATCCTACAAG TCCGTTGCGC ACACGCTTCG TATACCACCT GTCA	44
(2) INFORMATION FOR SEQ ID NO:61:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 33 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:61:	
GATCTGACAG GTGGTATACG AAGCGTGTGC GCA	33
(2) INFORMATION FOR SEQ ID NO:62:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 60 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:62: GATCTGTGTA AGAAGTGTGA TCCAACAGAG GTAGAGCTGG ACAATCAGAT AGTCACTGCA	60
(2) INFORMATION FOR SEQ ID NO:63:	

ff ff

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:	
GATCTGTGTA AGAAGGATGA GGACAGCGCT ACAGAAACCT GCTG	44
(2) INFORMATION FOR SEQ ID NO:64:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 44 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:	
AATTCAGCAG GTTTCTGTAG CGCTGTCCTC ATCCTTCTTA CACA	44
(2) INFORMATION FOR SEQ ID NO:65:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 62 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:	
GATCTGTGTA AGAAGGATGA GGACAGCGCT ACAGAAACCT GCTACGAGAA GGATGAGCTG	60
TG	62
(2) INFORMATION FOR SEQ ID NO:66:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 62 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:	
AATTCACAGC TCATCCTTCG CGTCGCAGGT TTCTGTAGCG CTGTCCTCAT CCTTCTTACA	60
CA	62

(2) INFORMATION FOR SEQ ID NO:67:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 59 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:67:  GATCTGTGTA AGAAGTCTGA TATCGATGAA GATTCCGCTA CAGAAACCTG CAGCACATG	59
(2) INFORMATION FOR SEQ ID NO:68:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 59 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:68:	
AATTCATGTG CTGCAGGTTT CTGTAGCGGA ATCTTCATCG ATATCAGACT TCTTACACA	59
(2) INFORMATION FOR SEQ ID NO:69:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 64 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:69:	
GATCTGTCTA AGAAGTCTGA TATCGATGAA GATTACAGAT TCTTCAGACT ATAGCTACTT	60
CTAA	64
(2) INFORMATION FOR SEQ ID NO:70:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs	

(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:70:	
AATCTTCATC GATATCAGAC TTCTTAGACA	30
(2) INFORMATION FOR SEQ ID NO:71:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 64 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:71:	
GATCTGGTTA AGAAGTCTGA TATCGATGAA GATTACCAAT TCTTCAGACT ATAGCTACTT	60
CTAA	64
(2) INFORMATION FOR SEQ ID NO:72:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 30 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:72:	
AATCTTCATC GATATCAGAC TTCTTAACCA	30
(2) INFORMATION FOR SEQ ID NO:73:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 41 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:73:	
ATTGTCCAGC TCTACCTCTG TTGGATCACA CTTCTTACAC A	41
(2) INFORMATION FOR SEQ ID NO:74:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 46 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:74:	
ACTCAAAGCA ACATTTGCGA TGAGGACAGC GCTACAGAAA CCTGCA	46
(2) INFORMATION FOR SEQ ID NO:75:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 57 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:75:	
GGTTTCTGTA GCGCTCTGCT CATCGCAAAT GTTGCTTTGA GTCGCAGTGA CTATCTG	57
(2) INFORMATION FOR SEQ ID NO:76:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 59 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:76:	

59

GCACCTACGA TAGGAACAAA TGCTACACGG CCGTGGTTCC GCTCGTGTAT GGTGGAGAG

(2) INFORMATION FOR SEQ ID NO:77:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 48 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:77:	
GAGCGGAACC ACGGCCGTGT AGCATTTGTT CCTATCGTAG GTGCTGCA	48
(2) INFORMATION FOR SEQ ID NO:78:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 50 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:78:  ACAAAAATGG TGGAAACTGC CCTTACGCCC GATGCATGCT ATCCGGACTG	50
(2) INFORMATION FOR SEQ ID NO:79:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 69 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:79:	
AATTCAGTCC GGATAGCATG CATCGGGCGT AAGGGCAGTT TCCACCATTT TTGTCTCTCC	60
ACCATACAC	69
(2) INFORMATION FOR SEQ ID NO:80:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 62 base pairs</li><li>(B) TYPE: nucleic acid</li></ul>	

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:80:	
ACAAAAATGG TGGAAACTGC CCTTACGCCC GATGCATGCT ATCCGGACAA GGATGAATTG	60
TG	62
(2) INFORMATION FOR SEQ ID NO:81:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 81 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:81:	
AATTCACAAT TCATCCTTGT CCGGATAGCA TGCATCGGGC GTAAGGGCAG TTTCCACCAT	60
TTTTGTCTCT CCACCATACA C	81
(2) INFORMATION FOR SEQ ID NO:82:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 88 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:82:	
GATCAGGTCG CTGCCATCCA AGACCCGAGG CTGTTCGCCG AAGAGAAGGC CGTCGCTGAC	60
TCCAAGTGCA AGTGTGCTCG TATTACTT	88
(2) INFORMATION FOR SEQ ID NO:83:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 88 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li></ul>	

(D) TOPOLOGY: linear

(xi)	SE	QUENCE	DESC	RIPTION: SE	EQ ID NO:83	:		
CTAGAAGT	AA	TACGAGC	ACA	CTTGCACTTG	GAGTCAGCGA	CGGCCTTCTC	TTCGGCGAAC	60
AGCCTCGG	GΤ	CTTGGAT	'GGC	AGCGACCT				88

- (2) INFORMATION FOR SEQ ID NO:84:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 10 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:84:

Cys Ala Ala Pro Lys Lys Lys Arg Lys Val 1 5 10

- (2) INFORMATION FOR SEQ ID NO:85:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 22 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:85:

Cys Ala Ala Lys Arg Pro Pro Ala Ala Ile Lys Lys Ala Ala Gly
1 5 10 15

Gln Ala Lys Lys Lys 20

- (2) INFORMATION FOR SEQ ID NO:86:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 4 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:86:	
His Asp Glu Leu 1	
(2) INFORMATION FOR SEQ ID NO:87:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 77 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:87:	
GCGATGACGA CGATAAGGCC CAAACGGAGA CCTGTACTGT TGCGCCTCGT GAACGGCAAA	60
ACTGCGGATT CCCGGAA	77
(2) INFORMATION FOR SEQ ID NO:88:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 66 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:88:	
GTTTTGCCGT TCACGAGGCG CAACAGTACA GGTCTCCGTT TGGGCCTTAT CGTCGTCATC	60
GCTTCA	66
(2) INFORMATION FOR SEQ ID NO:89:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 72 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:89:							
GTAACACCCT CTCAGTGCGC TAATAAAGGC TGCTGTTTTG ATGACACGGT ACGGGGCGTT	60						
CCGTGGTGCT TC							
(2) INFORMATION FOR SEQ ID NO:90:							
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 72 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>							
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:90:							
GCCCCGTACC GTGTCATCAA AACAGCAGCC TTTATTAGCG CACTGAGAGG GTGTTACTTC							
CGGGAATCCG CA							
(2) INFORMATION FOR SEQ ID NO:91:							
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 49 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>							
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:91:							
TACCCCAATA CAATTGACGT TCCGCCTGAA GAAGAGTGCG AGCCGTAAG	49						
(2) INFORMATION FOR SEQ ID NO:92:							
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 68 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>							

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:92:										
AATT	CTTAC	CG	GCTCGC	ACTC	TTCTTC	AGGC	GGCAAGTCAA	TTGTATTGGG	GTAGAAGCAC	60	
CACG	GAAC									68	

- (2) INFORMATION FOR SEQ ID NO:93:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 13 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:93:

Val Ala Val Gln Ser Ala Gly Thr Pro Ala Ser Gly Ser 1  $\phantom{\bigg|}$  5  $\phantom{\bigg|}$  10